



Figure 1

Sequence Name:

GW.S.ctg16335-000003.31.0

Figure 1A SEQ ID NO:01

LLAPTGSLFRNCTQDGWSETFPRPNLACGVNVNDSSNEKRSYLLKLKVMTVGYSSSLVM
LLVALGILCAFRLHCTRNYIHMHLFVFSFILRALSNFIKDAVLFSSDDVTYCDAHRGCKL
VMVLFXYCIMANYSWLVEGSTFThxLAISFFSERKYLQGFVAFWGWSPAIFVALWAIAR
HFLLEDVGCWDINANASIWWIIRGPVILSILNFINILRILMRKLRTQETRGNEVSHYK
RLARSTLLIPLFGIHYIVFAFSPEDAMEIQLFF

Figure 1B SEQ ID NO:02

CTCTTGGCACCCACAGGTTCCCTGTCGAAACTGCACACAGGATGGCTGGTCAGAAACC
TTCCCCAGGCCTAATCTGGCCTGTCGTTAATGTGAACGACTCTTCCAACGAGAAGCGG
CACTCCTACCTGCTGAAGCTGAAAGTCATGTACACCGTGGCTACAGCTCCTCCCTGGTC
ATGCTCCTGGTCGCCCTGGCATCCTCTGTGCTTCCGGAGGCTCCACTGCACTCGAAC
TACATCCACATGCACCTGTTCTGTCCTCATCCTCGTGCCTGTCCAACCTCATCAAG
GACGCCGTGCTCTCTCAGATGATGTCACCTACTGCGATGCCACAGGGCGGGCTGC
AAGCTGGTCATGGTGTGTT

Figure 1C SEQ ID NO:03

TACTGCATCATGGCCAACTAACACTCTGGCTGCTGGTGAAGGCTACACATNTC
CTCGCCATCTCCTTCTCTGAAAGAAAGTACCTCCAGGGATTGTGGCATTGGATGG
GGTTCTCCAGCCATTGGTGTGGCTATTGCCAGACACTTCTGGAAGATGTT
GGGTGCTGGGACATCAATGCCAACGCATCCATCTGGTGGATCATTGTCCTGTGATC
CTCTCCATCCTGATTAATTTCATCCTTCAAAACATTCTAAGAATCCTGATGAGAAA
CTTACAACCAAGAAACAAGAGGAAATGAAGTCAGCATTATAAGCGCCTGGCCAGGTCC
ACTCTCCTGCTGATCCCCCTTTGGCATCCACTACATCGTCTCGCCTCTCCCCAGAG
GACGCTATGGAGATCCAGCTGTTTTT

Figure 1D SEQ ID NO:04

CTCTTGGCACCCACAGGTTCCCTGTTCCGAAACTGCACACAGGATGGCTGGTCAGAAACC
TTCCCCAGGCCTAATCTGGCCTGTCGTTAATGTGAACGACTCTTCCAACGAGAAGCGG
CACTCCTACCTGCTGAAGCTGAAAGTCATGTACACCGTGGCTACAGCTCCTCCCTGGTC
ATGCTCCTGGTCGCCCTGGCATCCTCTGTGCTTCCGGAGGCTCCACTGCACTCGAAC
TACATCCACATGCACCTGTTCTGTCCTCATCCTCGTGCCTGTCCAACCTCATCAAG
GACGCCGTGCTCTCTCAGATGATGTCACCTACTGCGATGCCACAGGGCGGGCTGC
AAGCTGGTCATGGTGTGTT
TACTGCATCATGGCCAACTAACACTCTGGCTGCTGGTGAAGGCTACACATNTC
CTCGCCATCTCCTTCTCTGAAAGAAAGTACCTCCAGGGATTGTGGCATTGGATGG
GGTTCTCCAGCCATTGGTGTGGCTATTGCCAGACACTTCTGGAAGATGTT
GGGTGCTGGGACATCAATGCCAACGCATCCATCTGGTGGATCATTGTCCTGTGATC
CTCTCCATCCTGATTAATTTCATCCTTCAAAACATTCTAAGAATCCTGATGAGAAA
CTTACAACCAAGAAACAAGAGGAAATGAAGTCAGCATTATAAGCGCCTGGCCAGGTCC
ACTCTCCTGCTGATCCCCCTTTGGCATCCACTACATCGTCTCGCCTCTCCCCAGAG
GACGCTATGGAGATCCAGCTGTTTTT



Figure 2

Sequence Name:

GW.S.ctg16490-000000.17.0

Figure 2A SEQ ID NO:05

PTFILFSFQPGDKRTKHICVYWEGSEGGHWSTEGCSHVHSNGSYTKCKCFHLSSFAVLVA
LAPKDPVLTVITQVGLTISLLCLFLAILTFLLCRPIQNTSTSLHLELSLCLFLAHLLFLT
GINRTEPELCSIAGLLHFLYLACFTWMLLEGHLFLTVRNLKVANYTSTGRFKKRFMYP
VGYGIPAVIIAVSAIVGPQNYGTFTHCWLKLDKGFIWSFMGPVAVIILNLVFYFQVLWIL
RSKLSSLNKEVSTIQDTRVMTFKAISQLFILGCSWGLGFFMVEEVGKTIGSIIAYSFTII
NTLQGVLLFVVHCLLNQRQVR

Figure 2B SEQ ID NO:06

CCCACTTCTACTATTCTTTCCAGCCTGGTGACAAGAGAACAAAACATATCTGTGTC
TACTGGGAGGGATCAGAGGGAGGCCACTGGTCCACGGAGGGCTGCTCTCATGTGCACAGC
AACGGTTCTTACACCAAATGCAAGTGCCTCCATCTGTCCAGCTTGCCGTCTCGTGGCT
CTTGGCCCCAAGGAGGACCCGTGCTGACCGTGTACACCCAGGTGGGCTGACCATCTCT
CTGCTGTGCCTCTTCCATCCTCACCTCCTCCTGTGCCGGCCATCCAGAACACC
AGCACCTCCCTCCATCTAGAGCTCCCTCTGCCTCTTCCGTGCCACCTCCTGTTCTG
ACGGGCATCAACAGAACAGACTGAGCCTGAGGTGCTGTGCTCCATCTGCAGGGCTGTCAC
TTCCTCTACCTGGCTTGCTTCACCTGGATGCTCCTGGAAAGGGCTGCACCTCTCACC
GTCAGGAACCTCAAGGTGGCCAACTAACCCAGCACGGCAGATTCAAGAAGAGGTTCATG
TACCCCTGTAGGCTACGGGATCCCAGCTGTGATTATTGCTGTGTCAGCAATAGTTGGACCC
CAGAATTATGGAACATTTACTCACTGTTGGCTCAAGCTGATAAAAGGATTCATCTGGAGC
TTCATGGGGCCAGTAGCAGTCATTCTTGATAAAACCTGGTGTCTACTTCCAAGTTCTG
TGGATTTGAGAACAACTTCTCCCTCAATAAGAACGTTCCACCATTCAGGACACC
AGAGTCATGACATTAAAGCCATTCTCAGCTATTATCCTGGGTGTTCTGGGGCCTT
GGTTTTTATGGTTGAAGAACAGTAGGGAAGACGATTGATCAATCATTGCATACTCATTC
ACCATCATCAACACCCCTCAGGGAGTGTGCTTTGTGGTACACTGTCTCCTTAATCGC
CAGGTAAGG



Figure 3

Sequence Name:

GW.S.ctg13100-000000.33.0

Figure 3A SEQ ID NO:07

QHSDAVHDLLLDVITWVGILLSLVCLLICIFTFCFFRGLQSDRNTIHKNLCISLFVAELL
FLIGINRTDQOPACAVFAALLHFFFЛАAFTWMFLEGVQLYIMLVEVFESEHSRRKYFYLVG
YGMPALIVAVSAAVDYSYGTDKVCWLRLDTYFIWSFIGPATLIIMNVIFLGIALYKMFH
HTAILKPESGCLDNIKLKINIPPIKSIYIYMYICMCV

Figure 3B SEQ ID NO:08

CAGCACAGTGATGCGGTCCATGACCTCCTCTGGATGTGATCACGTGGTTGGAATTTG
CTGTCCTTGTTGTCTCCTGATTGCATCTCACATTGCTTTCCGGGGCTCCAG
AGTGACCGTAACACCATCCACAAGAACCTCTGCATCAGTCTTTGTAGCAGAGCTGCTC
TTCCTGATTGGGATCAACCGAACCTGACCAACCAATTGCCTGTGCTGTTTCGCTGCCCTG
TTACATTCTTCTTCTGGCTGCCTCACCTGGATGTTCTGGAGGGGGTGCAGCTTAT
ATCATGCTGGTGGAGGTTTGAGAGTGAACATTACGTAGGAAATACTTTATCTGGTC
GGCTATGGGATGCCTGCACTCATGGCTGTGTCAGCTGCAGTAGACTACAGGAGTTAT
GGAACAGATAAAAGTATGTTGGCTCGACTTGACACCTACTTCATTTGGAGTTATAGGA
CCAGCAACTTGATAATTATGCTTAATGTAATCTCCTGGATGCTTATATAAAATG
TTTCATCATACTGCTATACTGAAACCTGAATCAGGCTGTCTGATAACATCAAGTTAAA
ATTAATATTCCAATTATAAAATCTATTATATGTATATGCATGTGTGTG



Figure 4

Sequence Name:

GW.A.ctg12444-000001.0.2

Figure 4A SEQ ID NO:09

GNVAVFVYYKSIGPLLSSDNFLKPQNYDNSEEERVISSVISVSMSSNPPTLYELEK
ITFTLSHRKTDRYRSLCAFWNYSPTDMNGWSSEGCELTYSNETHTSCRCNHLTHFAILM
SSG PSI IKDYNILTRITQLGIIISLICLAI CIFT FWFFSEI QSTRTIHKNLCCSLFLAE
LVFLVGINTNTNKFCSI IAGLLHYFFLAFAWM CIEGIHLYLIVVGVIYNKGFLHKNFYI
FGYLSPAVVVGFSALGYRYGTTKVCWLSTENNFIWSFIGPACLIILVCIYKIVITI QK
SDDH

Figure 4B SEQ ID NO:10

GGCAATGTTGCAGTTGCATTGTATATTATAAGAGTATTGGTCCTTGCTTCATCATCT
GACAACCTCTTATTGAAACCTCAAATTATGATAATTCTGAAGAGGAGGAAAGAGTCATA
TCTTCAGTAATTCTCAGTCTCAATGAGCTAAACCCACCCACATTATATGAACTTGAAAAA
ATAACATTACATTAAGTCATCGAAAGGTACAGATAGGTATAGGAGTCTATGTGCATT
TGGAATTACTCACCTGATACCATGAATGGCAGCTGGTCTTCAGAGGGCTGTGAGCTGACA
TACTCAAATGAGACCCACACCTCATGCCGCTGTAATCACCTGACACATTGCAATTG
ATGTCCTCTGGTCCTTCATTGGTATTAAAGATTATAATTCTTACAAGGATCACTCAA
CTAGGAATAATTATTCACTGATTGTCTTGCATATGCATTTCACCTCTGGTTCTTC
AGTGAATTCAAAGCACCAGGACAACAATTCAACAAAATCTTGCTGTAGCCATTCTTCTT
GCTGAACCTGTTCTGTTGGGATCAATACAATACTAATAAGCTCTGTTCAATC
ATTGCCGGACTGCTACACTACTTCTTTAGCTGCTTGCATGGATGTGCATTGAAGGC
ATACATCTCTATCTCATGTTGTGGGTGTCACTACAACAAGGGATTTCACAGCAAGAAT
TTTATATCTTGGCTATCTAACGCCAGCCGTGGTAGTTGGATTTCGGCAGCACTAGGA
TACAGATATTATGGCACACAAAGTATGTTGGCTTAGCACCAGAAACAATTTATTGG
AGTTTATAGGACCAGCATGCCTAATCATTCTGTATGTATATAAAATTGTTATTACA
ATTCAAAAAAGTGTGATGATCAT



Figure 5

Sequence Name:

GW.S.ctg12789-000004.100.0

Figure 5A SEQ ID NO:11

GAWATTGCSVAALYLDSTACFCNHSTSFAILQIYEVQGPEEESLLRTLSFVGCGVSFCA
LTTFLLFLVAGVPKSERTTVHKNLTFSLASAEGFLMTSEWAKANEACVAVTVAMHFLFL
VAFSWMLVEGLLWRKVVAVSMHPGPGMRLYHATGWGVPGIVAVTLAMPHDYVAPGHC
WLNVHTNAIWAFVGPVLVLTVS

Figure 5B SEQ ID NO:12

GGTGCCTGGCCACACAGGCTGCTCCGTGGCTGCCCTGTACCTGGACTCCACCGCCTGC
TTCTGCAACCACAGCACCAGCTTGCCATCCTGCTGCAAATCTATGAAGTACAGAGAGGC
CCTGAGGAGGAGTCGCTGCTGAGGACTCTGTCAATTGTGGGCTGTGGCGTGCCTCTGC
GCCCTCACCAACACCTCTTGTCTTCCCTGGTGGCCGGGTCCCCAAGTCAGAGCGAAC
ACAGTCCACAAAGAACCTCACCTCTCCCTGCCCTCTGCCGAGGGCTTCCTCATGACCAGC
GAGTGGGCCAAGGCCAATGAGGTGGCATGTGTGGCTGTCACAGTCGCAATGCACTTCCTC
TTTCTGGTGGCATTCTCCTGGATGCTGGTGGAGGGGCTGCTGCTGTGGAGGAAGGTGGTA
GCTGTGAGCATGCACCCGGGCCAGGCATGCGGCTCTACCACGCCACAGGCTGGGCGTG
CCTGTGGGCATCGTGGCGGTACCCCTGGCATGCTCCCCATGACTACGTGGCCCCCGGA
CATTGCTGGCTCAATGTGCACACAAATGCCATCTGGGCCTTCGTGGGCCTGTGCTCTTC
GTGCTGACTGTGAGC



Figure 6

Sequence Name:

GW.A.ctg12776-000000.33.0

Figure 6A SEQ ID NO:13

MHSPRRTTLCMFIVIYSSKAALNWNYESTIHPLLHEHEPAGEEALRQKRAVATKSPTAE
EYTVNIEISFENASFLDPIKAYLNSLSFPIHGNNTDQITDILSINVTTVCRPAGNEIWCS
CETGYGWPRERCLHNLICQERDVFLPGHHCSCLKELPPNGFCLLQEDVTLNMRVRLNVE
FQEDLMNTSSALYRSYKTDLETARKGYGILPGFKGVTVTGFKSGSVVVTYEVKTPPSLE
LIHKANEQVQSLNQTYKMDYNSFQAVTINESNFFVTPEIIFEGDTVSLVCEKEVLSSNV
SWRYEEQQLEIQNNSRFSIYTALFNNMTSVSKLTIHNITPGDAGEYVCKLILDIYECK
KKIDVMPIQILANEEMKVMCDNNPVSNCQGNVNWSKVEWKQEGKINIPGTPETDIDS
SCSRYTLKADGTQCPGSSGTTVIYTCFISAYGARGSANIKVTFISVANLTITPDPISV
SEGQNFISIKCISDVSNYDEVYWNTSAGIKIYQRFYTTRRYLDGAESVLTVKTSTREWNGT
YHCIFRYKNSYSIATKDVIVHPLPLKLNIMVDPLEATVSCSGSHHIKCCIEEDGDYKVTF
HTGSSSLPAAKEVNKKQVCYKHNFNASSVSWCSKTVDCCHFTNAANNSVWSPSMKNLV
PGENITCQDPVIGVGEPGKVIQKLCRFSNVPSSPESPIGGTITYKCVGSQWEEKRNDIS
APINSLLQMAKLIKSPSQDEMLPTYLKDLSISIDKAEHEISSSPGSLGAIINILDLLSTV
FTQVNSEMMTBLSTVNVLGKPVLNTWKVLQQQWTNQSSQLLHSVERFSQALQSGDSPPL
SFSQTNVQMSMVIKSSHPTYQQRFVFPYFDLWGNVVIDKSYLENLQSDSSIVTMAFPT
LQAILAQDIQENNFAESLVMTTVSHNTTMRISMTFKNNSPSGGETKCVFWNFRANN
TGGWDSSGCYVEEGDGDNVTCICDHLTFSILMSPDSDPSSLGILLDIISYVGVGFSI
LSLAACLVVEAVVWKSVTKNRTSYMRHTCIVNIAASLLVANTWFIVVAAIQDNRYILCKT
ACVAATFFIHFYLSVFFWMLTLGLMLFYRLVFLHETSRSRSTQKIAFCLGYGCPLAISV
ITLGATQPREVYTRKNVCWLWEDTKALLAFAIPALIIVVVNITITIVVITKILRPSIGD
KPKQEKSSLFQISKSIGVLTPLLGLTWGFLTTVFPGTNLVHIIFAILNVFQLFILLF
GCLWDLKQEALLNKFSLSRWSSQHSKTSLGSTPVFSMSSPISRRFNNLFGKTGTYNVST
PEATSSLENSSSASSLLN

Figure 6B SEQ ID NO:14

ATGAAATCCCCAAGGAGAACCACTTGTGCCTCATGTTATTGTGATTTATTCTTCCAAA
GCTGCACTGAACACTGGAATTACGAGTCTACTATTACATCCTTGAGTCATGAACATGAA
CCAGCTGGTGAAGAGGCACTGAGGCCAAAACGAGCCGTTGCCACAAAAGTCCTACGGCT
GAAGAATAACACTGTTAATATTGAGATCAGTTGAAAATGCATCCTCCTGGATCCTATC
AAAGCCTACTTGAAACAGCCTCAGTTCCATTACATGGGAATAACACTGACCAAATTAC
GACATTGAGCATAAAATGTGACAACAGTCAGCAGACCTGCTGGAAATGAAATCTGGTGC
TCCTGCAGACAGGTTATGGGTGCCCTCGGAAAGGTGTCTTACAATCTCATTGCAA
GAGCGTGACGTCTCCCTCCCAGGGCACCATTCAGTTGCCTTAAAGAACTGCCTCCCAAT
GGACCTTTGCCTGCTTCAGGAAGATGTTACCTGAACATGAGAGTCAGACTAAATGTA
GGCTTCAAGAAGACCTCATGAACACTCCCTCCGCCCTCTATAGGTCTACAAGACCGAC
TTGGAAACAGCGTCCGGAAGGGTACGGAATTACAGGCTCAAGGGCGTGACTGTG
ACAGGGTTCAAGTCTGGAAGTGTGGTTGTGACATATGAAGTCAAGACTACACCACATCA
CTTGAGTTAACATAAGCCAATGAACAAGTTGTACAGAGCCTCAATCAGACCTACAAA
ATGGACTACAACCTTCAAGCAGTTACTATCAATGAAAGCAATTCTTGTACACACCA
GAAATCATCTTGAAGGGACACAGTCAGTCTGGTGTGAAAGGAAGTGTCTCC
AATGTGTCTTGGCGCTATGAAGAACAGCAGTTGGAATCCAGAACAGCAGCAGATTCTCG
ATTACACCGCACTTCAACACATGACTCGGTGCCAAGCCTCACCATCCACAACATC
ACTCCAGGTGATGCAGGTGAATATGTTGCAAACACTGATATTAGACATTTGAATATGAG
TGCAAGAAGAAAATAGATGTTATGCCATCAAATTGGAATGAAGAAATGAAGGTG
ATGTGCGACAACAATCCTGTATTTGAACTGCTGCAGTCAGGTAATGTAATTGGAGC
AAGTAGAATGGAAGCAGGAAGGAAAATAAATATTCCAGGAACCCCTGAGACAGACATA
GATTCTAGCTGCAGCAGATAACCCCTCAAGGCTGATGGAACCCAGTGCCAAGCGGGTCG

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TCTGGAACACAGTCATCTACACTTGTGAGTCATCAGTGCCTATGGAGCCAGGGCAGT
GCAAACATAAAAGTGACATTCTCATCTGTGGCAATCTAACATAACCCCGGACCCAATT
TCTGTTCTGAGGGACAAAACCTTCTATAAAATGCACTCAGTGATGTGAGTAACATATGAT
GAGGTTATTGGAACACTCTGCTGGATTAAAATACCAAAGATTATACCGAGG
AGGTATCTTGATGGAGCAGAATCAGTACTGACAGTCAGACCTCGACCAGGGAGTGGAAAT
GGAACCTATCACTGCATATTAGATAAGAATTCACTACAGTATTGCAACCAAAGACGTC
ATTGTTACCCGCTGCCTCTAAAGCTGAACATCATGGTGATCCTTGGAAAGCTACTGTT
TCATGCAGTGGTCCCACATCACATCAAGTGCTGCATAGAGGAGGATGGAGACTACAAAGTT
ACTTCCATACGGGTTCCCTCATCCCTCCTGCTGCAAAGAAGTTAACAAAAAACAAAGTG
TGCTACAAACACAATTCAATGCAAGCTCAGTTCCCTGGTGTCAAAAACACTGTTGATGTG
TGTGTCACTTACCAATGCTGCTAATAATTCACTGCTGGAGCCATCTATGAAGCTGAAT
CTGGTCTGGGAAACATCACATGCCAGGATCCCGTAATAGGTGTCGGAGAGCCGGGG
AAAGTCATCCAGAAGCTATGCCGGTTCTCAAACGTTCCAGCAGCCCTGAGAGTCCCATT
GGCAGGACCATCACTTACAAATGTTAGGCTCCAGTGGGAGGAGAAGAGAAATGACTGC
ATCTCTGCCCAATAAACAGTCTGCTCCAGATGGCTAAGGCTTGTGATCAAGAGCCCTCT
CAGGATGAGATGCTCCCTACATACCTGAAGGATCTTCTATTAGCATAGACAAAGCGGAA
CATGAAATCAGCTTCTCTGGGAGTCTGGGAGCATTATTAACATCCTGATCTGCTC
TCAACAGTCCAACCCAAGTAAATTCAAGAATGATGACGCACGTGCTCTACGGTTAAT
GTCATCCTGGCAAGCCGCTTGAACACCTGAAAGGTTTACAACAGCAATGGACCAAT
CAGAGTTCAAGCTACTACATTCACTGGAAAGGATTTCCAAGCATTACAGTCGGAGAT
AGCCCTCCTTGTCTCCAAACTAATGTCAGATGAGCAGCATGGTAATCAAGTCC
AGCCACCCAGAAACCTATCAACAGAGGTTGTTCCACTTTGACCTCTGGGCAAT
GTGGTCATTGACAAGAGCTATCTAGAAAACCTGAGTCGGATTGCTATTGTCACCATG
GCTTCCCAACTCTCCAAGCCATCCTGGCCAGGATATCCAGGAAAATAACTTGCAGAG
AGCTTAGTGTGACAAACCCTGTCAGCCACAATACAACATGCCATTAGGATTCAATG
ACTTTAAAGAACAAATAGCCCTCAGGGCGAACAGAAGTGTGTTCTGGAAACTTCAGG
CTGCCAACACAGGGGGTGGACAGCAGTGGGTGCTATGTAGAAGAAGGTGATGG
GACAATGTCACCTGTATCTGTGACCACCTAACATCATCTCCATCCTCATGCTCC
TCCCCAGATCCTAGTTCTCTCTGGGAAACTCCTGGATATTATTCTTGTGTTCTGGG
GGCTTTCCATCTGAGCTTGGCAGCCTGCTAGTTGTGGAAGCTGTGGTGTGAAATCG
GTGACCAAGAACCGGACTCTTATATGCGCCACACCTGCATAGTGAATATCGCTGCCTCC
CTCTGGTCGCAACACCTGGTTCATTGTGGTCGCTGCCATCCAGGACAATCGCTACATA
CTCTGCAAGACAGCCTGTGTGGCTGCCACCTTCTCATCCACTTCTTACCTCAGCGTC
TTCTCTGGATGCTGACACTGGGCTCATGCTGTTCTATGCCCTGGTTTCATTCTGCAT
GAAACAAGCAGGTCCACTCAGAAAGCCATTGCCCTCTGCTTGGCTATGGCTGCCACTT
GCCATCTCGTCATCACGCTGGAGCCACCCAGCCCCGGAAAGTCTATACGAGGAAGAAT
GCTGTTGGCTCAACTGGGAGGACACCAAGCCCTGCTGGCTTGCCTACCCAGCACTG
ATCATTGTGGTGGTGAACATAACCATCACTATTGTGGTCATACCAAGATCCTGAGGCT
TCCATTGGAGACAAGCCATGCAAGCAGGAGAAGAGCAGCCTGTTCACTCAGCAAGAGC
ATTGGGGTCTCACACCACTCTGGGCTCACTGGGTTTGGTCTCACCACGTGTT
CCAGGGACCAACCTTGTGTTCCATATCATATTGCCATCCTCAATGTCTCCAGGGATTA
TTTCATTGTCGAGATGGTCTCACAGCACTAAAGCTAACATCCCTGGGTTCATCCACA
CCTGTGTTCTATGAGTTCTCAACATCAAGGAGATTAAACAATTGTTGGTAAACAA
GGAACGTATAATGTTCCACCCAGAAGCAACCAGCTCATCCCTGGAAAACACTCATCCAGT
GCTTCTCGTTGCTCAAC



Figure 7

Sequence Name:

GW.S.ctg12776-000000.175.0

Figure 7A SEQ ID NO:15

ILNSKSISNWTIFIRDRNSSYILLHSVNSFARRLFIDNIPVDISDVFIHTMGTISGDNIG
KNFTFSMRINDTSNEVTGRVLISRDELKVPSPSQVISIAFPTIGAILEASLLENVTNG
LVLSAILPKELKRISLIFEKISKSEERRTQCVGWHHSVENRWDQQACKMIQENSQQAVCKC
RPSKLFTSFSILMSPHILESILTYITYVGLGISICSLILCLSIEVLVWSQVTKTEITYL
RHVCIVNIAATLLMADVWFIVASFSLSGPITHHKGCVAAFFVHFFYLSVFFWMLAKALLI
LYGIMIVFHTLPKSVLVASLFSVGYGCPLAIAAITVAATEPGKGYLRPEICWLWDMTKA
LLAFVIPALAIVVNLITVTLVIVKTQRAAIGNSMFQEVRRAIRISKNIAILPLLGLTW
GFGVATVIDDRSLAFHIIFSLNAFQFFILVFGTILDPKV

Figure 7B SEQ ID NO:16

ATTCTAACAGAAAAGCATCTCAACTGGACTTTCATTGTGACAGAACAGCAGCTAT
ATCCTGCTACATTCACTCCTTGCAAGAAGGCTATTAGATAACATCCCTGTT
GACATATCAGATGTCTCATTCAACTATGGGACCAACCATACTGGAGATAACATTGGA
AAAAATTCTACTTTCTATGAGAATTAAATGACACCAGCAATGAAGTCACTGGGAGAGTG
TTGATCAGCAGAGATGAACCTCGGAAGGTGCCTCCCTCTCAGGTCACTCAGCATTGCA
TTCCAACATTGGGCTATTGGAAAGCCAGTCTTGGAAAATGTTACTGTAAATGGG
CTTGTCTGTCTGCCATTGCCAAGGAACCTAAAAGAATCTCACTGATTGAAAAG
ATCAGCAAGTCAGAGGAGAGGACACAGTGTGTTGGCTGGCACTCTGTGGAGAACAGA
TGGGACCAGCAGGCCTGCAAATGATTCAAGAAAATCCCAGCAAGCTGTTGCAAATGT
AGGCCAAGCAAATTGTTACCTCTCAATTCTATGTCACCTCACATCTTAGAGAGT
CTGATTCTGACTTACATCACATATGTAGGCTGGCATTCTATTGAGCTGATCCT
TGCTGTCCATTGAGGTCTAGTCTGGAGCCAAGTGACAAAGACAGAGATCACCTATT
CGGCATGTGTGCATTGTTAACATTGAGCCACTTGTGATGGCAGATGTGTTGTCATT
GTGGCTCCTTCTTAGTGGCCAATAACACACCAAGGGATGTGTTGAGCCACATT
TTGTTCATTCCTTACCTTCTGTATTGTTCTGGATGCTGGCCAAGGCACCTTATC
CTCTATGGAATCATGATTGTTCCATACCTTGGCCAAGTCAGTCTGGCATCTCTG
TTTCAGTGGCTATGGATGCCCTTGGCATTGCTGCCATCACTGTTGCTGCCACTGAA
CCTGGCAAAGGCTATCAGACCTGAGATCTGCTGGCTCAACTGGGACATGACCAAGGCC
CTCCTGGCCTCGTGATCCCAGCTTGGCCATCGTGGTAGTAAACCTGATCACAGTCACA
CTGGTATTGTCAAGACCCAGCGAGCTGCCATTGCAATTCCATGTTCCAGGAAGTGAGA
GCCATTGTGAGAATCAGCAAGAACATGCCATCCTCACACCACTCTGGACTGACCTGG
GGATTGGAGTAGCCACTGTCATGATGACAGATCCCTGGCCTCCACATTATCTTCTCC
CTGCTCAATGCATTCCAGGGTTCTCATCCTAGTGTGTTGGAACCACCTGGATCCAAAG
GTA



Figure 8

Sequence Name:

GW.S.ctg16790-000000.13.0

Figure 8A SEQ ID NO:17

GTTGDWSSEGCSTEVRPEGTVCCDHLTFALLPTLDQSTVHILTRISQAGCGVSMIFL
AFTIILYAFRLRLSRERFKSEDAPKIHVALGGSLFLLNLAFLVNVGSGSKGSDAACWARGA
VFHYFLLCAFTWMGLEAFHLYLLAVRVFNTYFGHYFLKLSLVGWLPAKMVGTSANSY
GLYTIRDRENRTSLELCWFREGTTMYALYITVHGYFLITFLFGMVVLALVVWKIFTLSRA
TAVKERGKNRKKVLTILGLSSLVGVTWGLAIFTPLGLSTVYIFALFNSLQVDFYILIFY

Figure 8B SEQ ID NO:18

GGGACCACTGGAGACTGGTCTTCTGAGGGCTGCTCACGGAGGTAGACCTGAGGGGACC
GTGTGCTGCTGTGACCACCTGACCTTTTCGCCCTGCTCCTGAGACCCACCTGGACCAG
TCCACGGTGATATCCTCACACGCATCTCCCAGGCGGGCTGTGGGTCTCCATGATCTTC
CTGGCCTTCACCAATTATTCTTATGCCTTCTGAGGCTTCCCGGGAGAGGTTCAAGTCA
GAAGATGCCCAAAGATCCACGTGGCCCTGGTGGCAGCCTGTCCTCCTGAATCTGGCC
TTCTGGTCAATGTGGGAGTGGCTAAAGGGTCTGATGCTGCCCTGCTGGGCCGGGG
GCTGTCTTCACTACTCCTGCTGTGCCCTCACCTGGATGGGCCCTGAAGCCTTCCAC
CTCTACCTGCTCGTGTCAAGGTCTTCAACACCTACTCGGGCACTACTTCTGAAGCTG
AGCCTGGTGGCTGGGCCCTGCCGCCCTGATGGTCATCGGCAGTGGAGTGCCAACAGC
TACGGCCTTACACCATCCGTGATAGGGAGAACCGCACCTCTGGAGCTATGCTGGTTC
CGTGAAGGGACAACCATGTACGCCCTATATCACCGTCCACGGCTACTTCTCATCACC
TTCTCTTGGCATGGTGGCTGGCCCTGGTGGTCTGGAAGATCTCACCCGTCCCCGT
GCTACAGCGGTCAAGGAGCGGGGGAGAACCGGAAGAAGGTGCTCACCCGTGGGCCCT
TCGAGCCTGGTGGGTGTGACATGGGGTTGCCATCTCACCCGTGGGCCCTCCACC
GTCTACATCTTGCACCTTCAACTCCTGCAAGTTGATTTACATATTGATCTTCTAT



Figure 9

Sequence Name:

GW.S.ctg12776-000000.172.0

Figure 9A SEQ ID NO:19

NHILD TAAIS NWAFI PNKNASS DLLQSVNL FARQL HHN SENIV NELFI QT KGF HINHN
TSEKSLNFSMSMNN TEDI LGMVQI PRQELRKL WPNA SQAISIA FPTLGAI LREAH LQNV
SLPRQVNGLVLSVVL PERI QEI ILT FEKINKTRNARA QCVGWHSKRRWDEKACQMMLDI
RNEVKCRCNYTSVMSFSILMSSKSMTDKVLDYITCIGLSVSILSLVLCLI EATVWSRV
VVTEISYMRHVCIVNIAVSLLTANVWFIIGSHFNIAQDYNMVCAVTFFSHFFYLSLFW
MLFKALLI IYGILVIFRRMMKS RMMVIGFAIGYGCPLIIAVTTVAITEPEKGYIRPEACW
LNWDNTKALXAFAI PAFVIVAVNLIVVAVNTQRPSIGSSKSQDVVIIMRISKVAIL
TPLLGLTWGFIATLIEGSLTFHIIIFALLNAFQFFILLFGTIMDHKV

Figure 9B SEQ ID NO:20

AACCACATCCTCGACACAGCAGCCATTCAA ACTGGGCTTCATTCCAACAAAAATGCC
AGCTCGGATTGTTGCAGTCAGTGAATTGTTGCCAGACA ACTCCACATCCACAATAAT
TCTGAGAACATTGTGAATGA ACTCTTCATTCA GACA AAAAGGGTTCACATCAACCATAAT
ACCTCAGAGAAAAGCCTCAATTCTCCATGAGCATGAACAATACCACAGAAGATATCTTA
GGAATGGTACAGATTCCCAGGCAAGAGCTAAGGAAGCTGTGGCCAAATGCATCCCAAGCC
ATTAGCATAGCTTCCCAACCTGGGGCTATCCTGAGAGAAGGCCACTTGCAAAATGTG
AGTCTCCCAGACAGGTAATGGCTGGTCTATCAGGGTTTACCA GAAAGGTTGCAA
GAAATCATACTCACCTCGAAAAGATCAATAAAACCCGCAATGCCAGAGCCCAGTGTGTT
GGCTGGCACTCCAAGAAAAGGAGATGGATGAGAAAGCGTGC CAAATGATGTTGGATATC
AGGAACGAAGTGAATGCCGCTGTA ACTACACCAGTGTGGT GATGTCTTTCCATTCTC
ATGTCCTCCAATCGATGACCGACAAGTTCTGGACTACATCACCTGCATTGGCTCAGC
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GCTGTAATCTGATTGGTTGGTGTGCTGCAACACTCAGAGGCCCTCTATTGGC
AGTTCCAAGTCTCAGGATGTGGTCATAATTATGAGGATCAGCAAAATGTGCCATC
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TTGGAAACCATTATGGATCACAAGGTA